FIGURE 1

Vκ Domain

```
            1         10        20        30        40
CD40    DIVLTQSPATLSVTPGDRVSLSCRASQSISDYLHWYQQKSHESPRLLIK
        |         || | ||                     ||||       |
VKIII   EIVLTQSPATLSLSPGERATLSCRASQSVSSYLAWYQQKPGQAPRLLIY
                                                        *

           50        60        70        80        90        100
CD40    YASHSISGIPSRFSGSGSGSDFTLSINSVEPEDVGIYYCQHGHSFPRTFGGGTKLEIK
                            |     | | |       |||        |
VKIII/JK4 DASNRATGIPARFSGSGSGTDFTLTISSLEPEDFAVYYCQQRSNWPLTFGGGTKVEIK
```

VH Domain

```
            1         10        20        30        40
CD40    QIQLVQSGPELKKPGETVRISCKASGYAFTTTGMQWVQEMPGKGLKWIG
         |        |       || ||          |          ||| | | |
VH7     QVQLVQSGSELKKPGASVKVSCKASGYTFTSYAMNWVRQAPGQGLEWMG
                   *                    *         **     *  *

           50        60        70        80 abc      90
CD40    WINTHSGVPKYVEDFKGRFAFSLETSANTAYLQISNLKNEDTATYFCVR

VH7     WINTNTGNPTYAQGFTGRFVFSLDTSVSTAYLQISSLKAEDTAVYYCAR
                                                       *
```

```
          abcde          110
CD40    SGNGNYDLAYFAYWGQGTLVTVSA
                            |
JH4     _____YFDYWGQGTLVTVSS
```
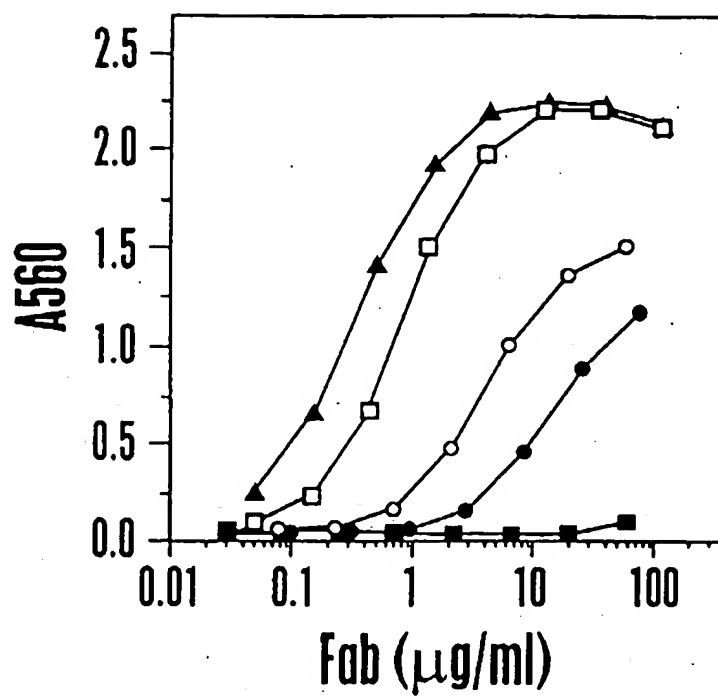
FIGURE 2

Fig. 2A

Fig. 2B

FIGURE 3

FIGURE 4

Figure 4A

Figure 4B

FIGURE 5

V_H Domain

```
                              FR1                    CDR-H1        FR2              CDR-H2                      FR3

                    1                   2            3             4                5          6               7          8          9
                    1234567890123456789012345       6789012345    6789012345       0123456789012345           6789012345678901 2abc345678901234

NMC4        VQLLESGPGLVAPSQSLSITCTVS              GFSLTDYGVD    WVRQPPGKGLEWLG     MIWGDGSTDYNSALKS           RLSITKDNSKSQVFLKMNSLQTDDTARYYCVR
VH3-72      EVQLVESGGGLVQPGGSLRLSCAAS             ----------    WVRQAPGKGLEWVG     ----------------           RFTISRDDSKNSLYLQMNSLKTEDTAVYYCAR


            CDR-H3
            1       1
            0       1
            567890abcdef12  3456789012
NMC4        DPADYGNYDYALDY  WGQGTSVTVS    (SEQ ID NO:5)
JH6a        --------------  WGQGTTVTVS    (SEQ ID NO:7)
```
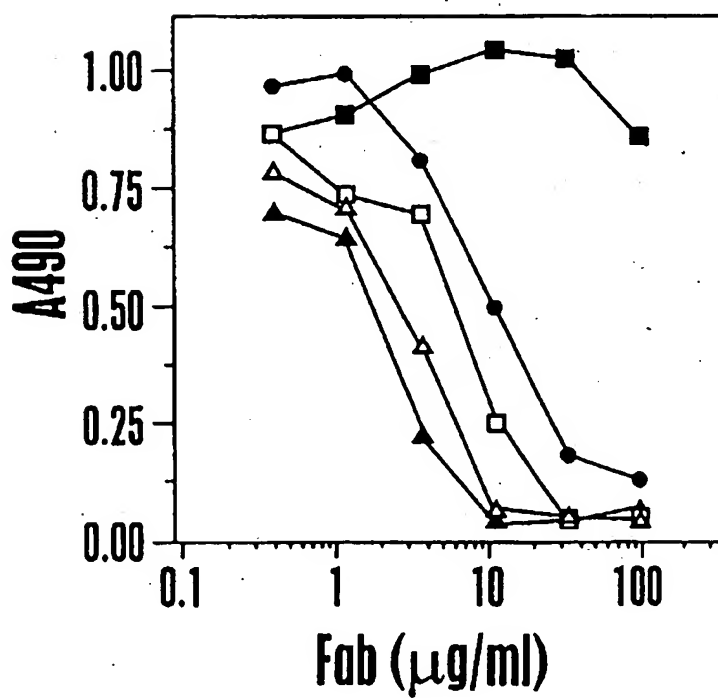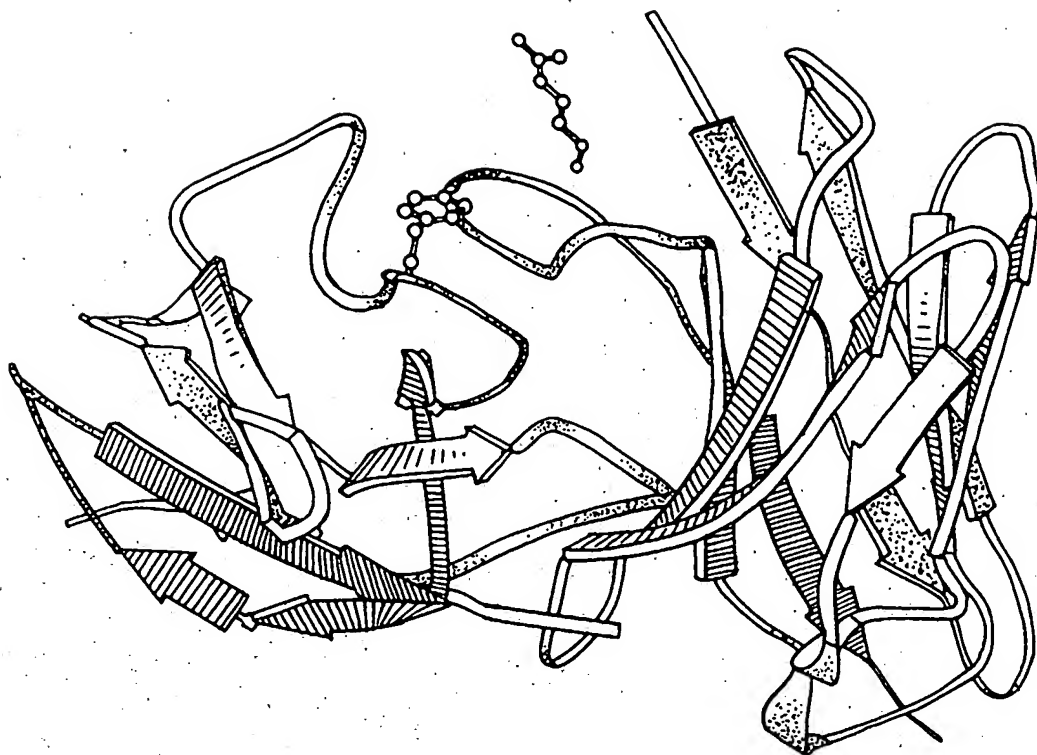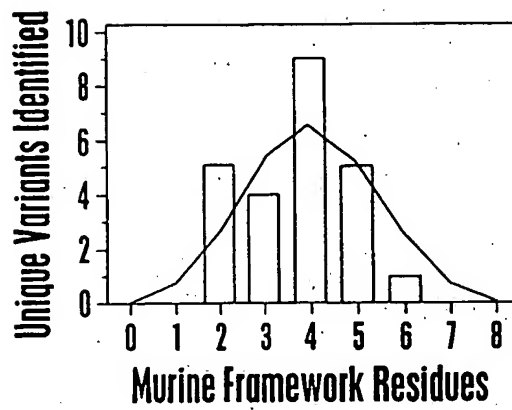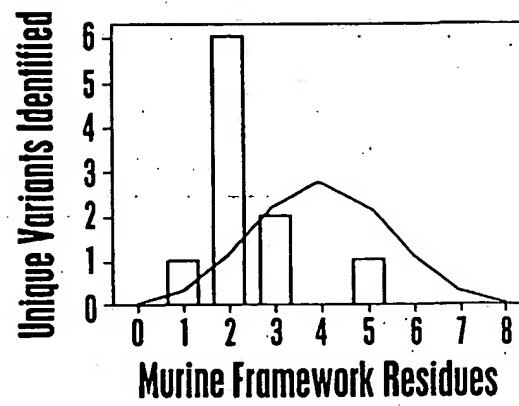
V_κ Domain

V_κ Domain

```
                         FR1                    CDR-L1         FR2              CDR-L2      FR3

                1          2                    3              4                5           6          7          8
                1234567890123456789 0123        45678901234    56789012345 6789 0123456     78901234567890123456789012345678

NMC4 VL         SSLSASLGDRVTISC                SASQDINKYLN    WYQQKPDGAVKLLIF   YTSSLHS     GVPSRFSGSGSGTDYSLTISNLEPEDIATYYC
VKIDPK9/012     DIQMTQSPSSLSASVGDRVTITC        -----------    WYQQKPGKAPKLLIY   -------     GVPSRFSGSGSGTDFTLTISSLQPEDFATYYC


            CDR-L3
            1
            9          0
            901234567  8901234567
NMC4 VL     QQYEKLPWT  FGGGTKLEVK    (SEQ ID NO:6)
JK4         ---------  FGGGTKVEIK    (SEQ ID NO:8)
```

FIGURE 6

vWF Elisa of Combinatorial Positives